

Enhancing Software-Related Information Extraction via Single-Choice Question Answering with Large Language Models

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Shared Task: Software Mention Detection in Scholarly Publications (SOMD)

- 1 Identify Mentions of 5 Software related entity types incl. Intention (e.g., **Application, Plugin, Operating System**)
- 2 Detect 10 attributive entities (e.g., **Version, License, Developer**)
- 3 Identify 13 typed relations between the Mentions (e.g., **version of, license of**)

Sentence:

"This example file runs on



Motivation

- Tracking Software Usage
- Foundation for Reproducibility Studies
- Foster Transparency in Research

Given Dataset

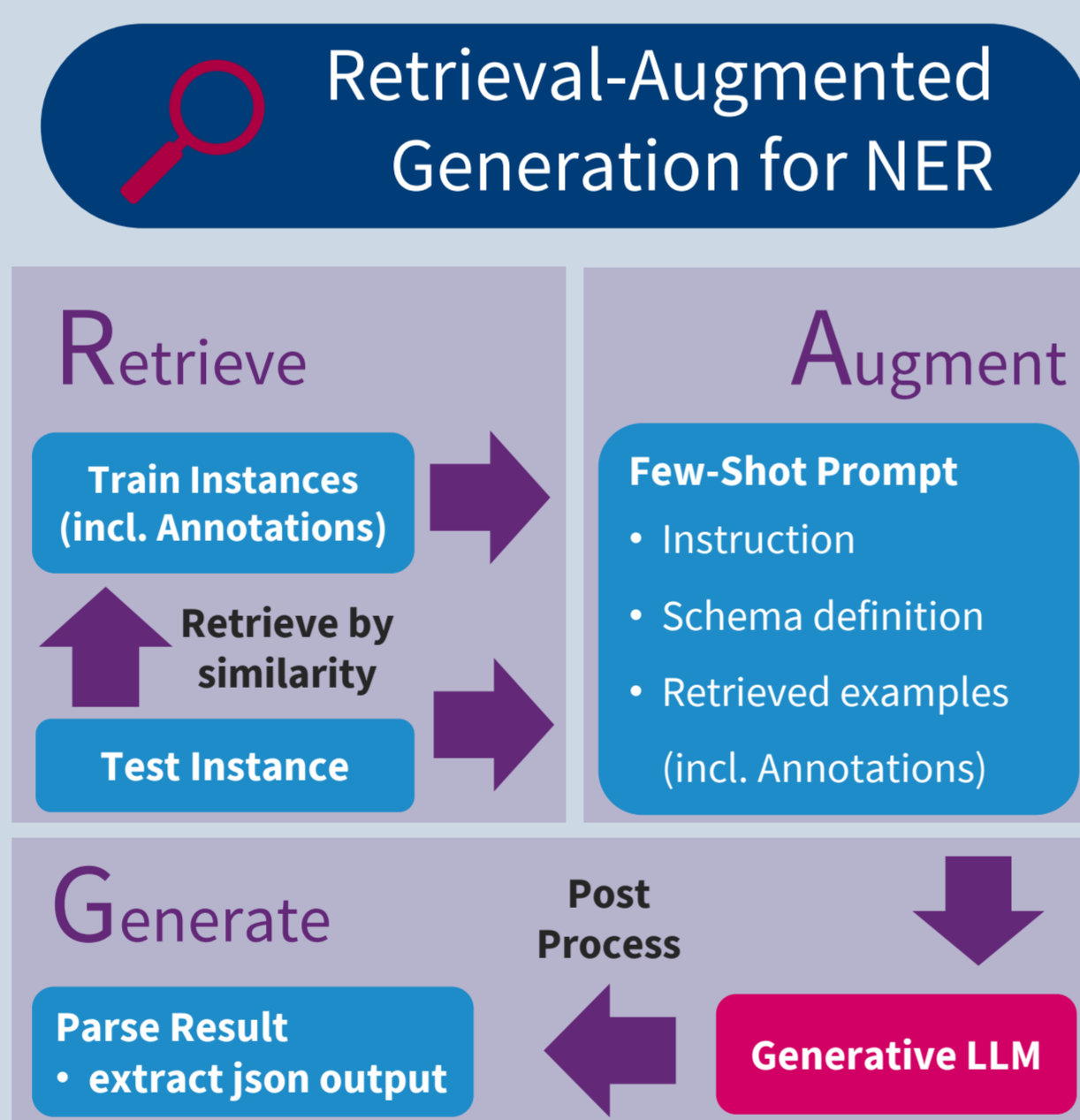
- ~1,400 Publication
- ~48,000 Sentences
- ~3,800 Software Mentions

- Bio and medical domain (Pubmed + PLoS)
- Based on SoMeSci
- used for SoftwareKG

		Train	Test		Test share
All sentences		39,769 (100.0%)	8181 (100.0%)	ST1 Data	17.1%
Incl. entity annotations	(ST1 Task)	2,326 (5.8%)	373 (4.6%)	ST2 Data	13.8%
Incl. attributive annotations	(ST2 Task)	1,690 (4.3%)	230 (2.8%)		12.0%
Incl. relation annotations	(ST3 Task)	1,075 (2.7%)	128 (1.6%)	ST3 Data	10.6%

Approach

- Show expected scope of task
- Show similar entities/relations
- Show similar contexts
- **Idea:** Use Training Contexts as examples



Sentence:

"This example file runs on **BEAST 2.1.0** with the **MultiTypeTree** package [..]."

Example Question:

Which of the statements about the Version '2.1.0' is true?

- '2.1.0' is the version of 'BEAST'.
- '2.1.0' is the version of 'MultiTypeTree'.

Derived Relation:



Relation Extraction as Question Answering

Observations:

- Every attributive entity has exactly one related entity
- If there is only one candidate entity in a sentence we know the answer (i.e., a "necessary" relation)
- For the others, we need to decide between a few (cf. Single Question Answering)
- For relations with multiple candidate, we reformulate the RE Task as a single-choice QA task

Results

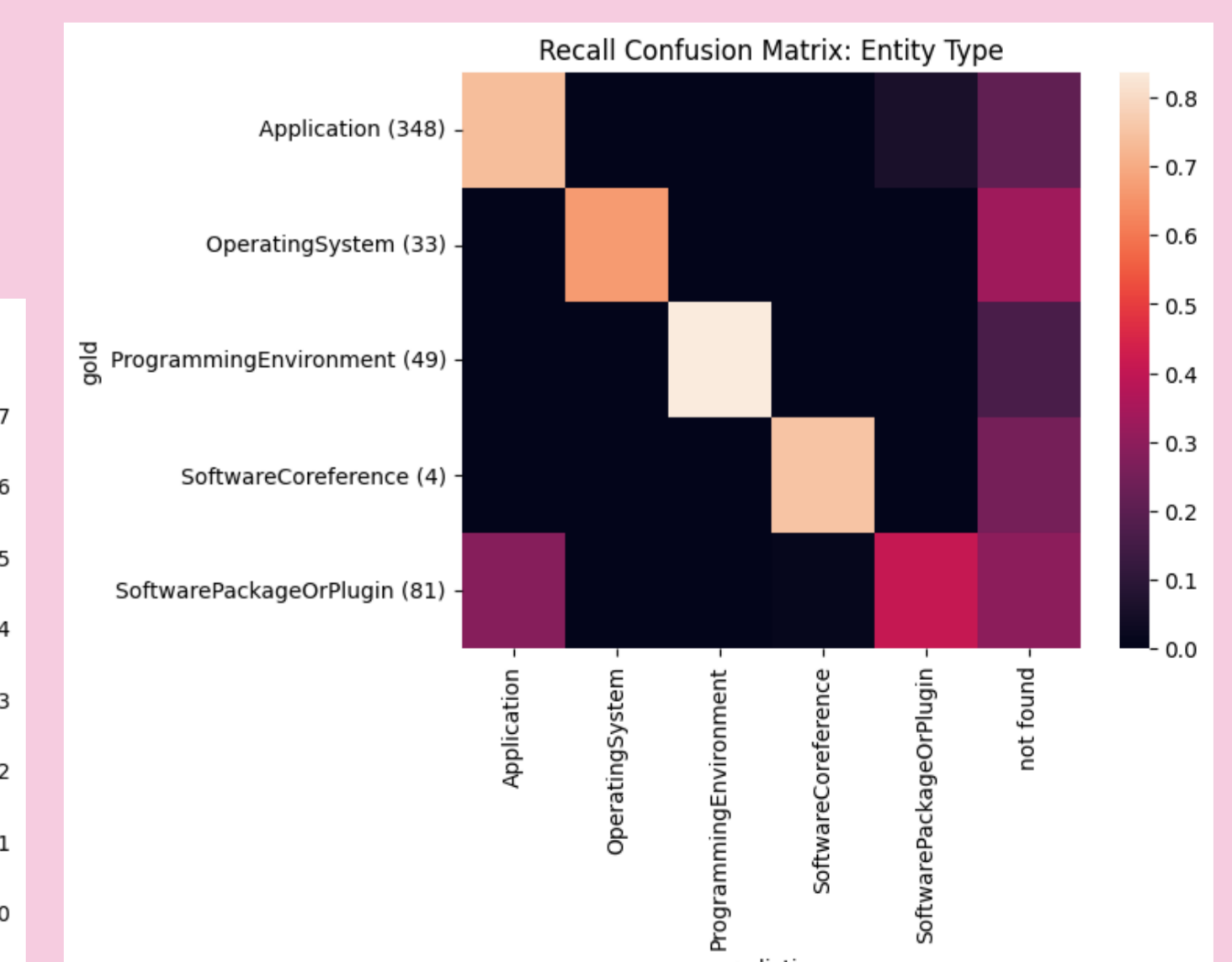
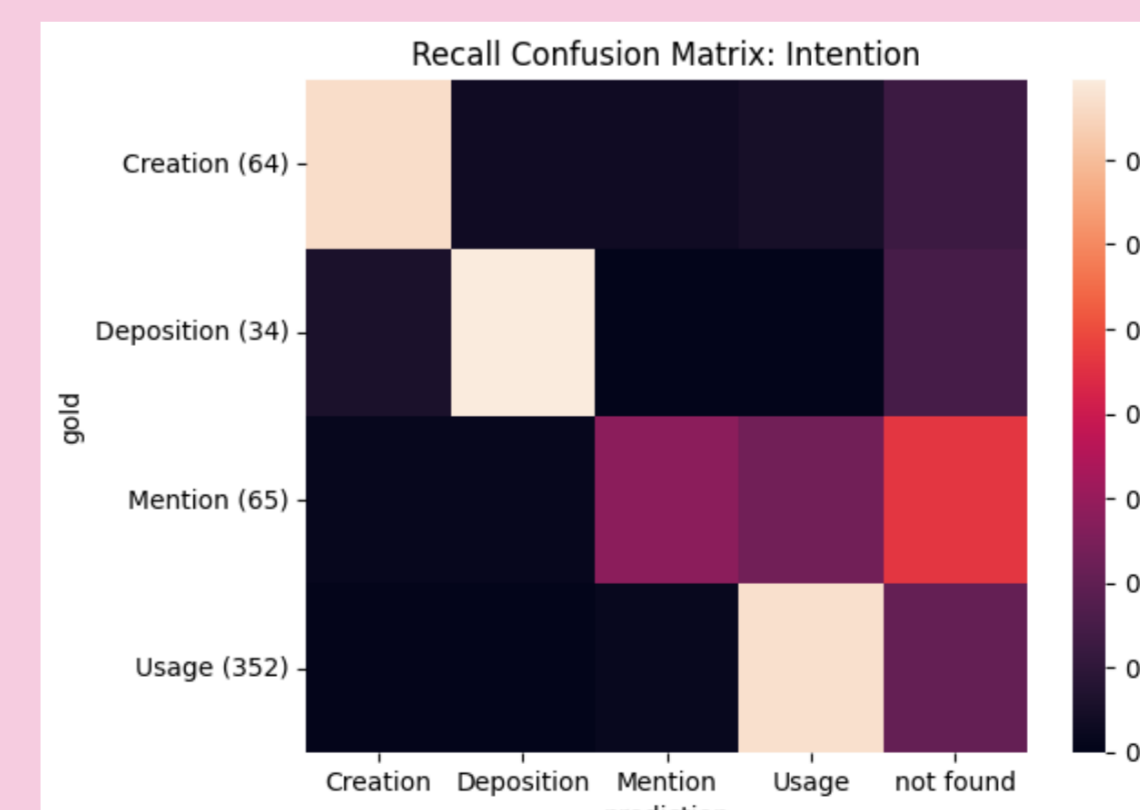
Retrieval Examples: Entity Similarity

entity	label	sim	split	sentence
PhosphOrtholog	Application	1.00	test	To this end , we have developed an automated web - based tool , PhosphOrtholog , which allows batch processing and mapping of large species - specific PTM datasets to compare overlap at a site - specific level .
SNPdetector	Application	0.93	train	We developed a software tool , SNPdetector , for automated identification of SNPs and mutations in fluorescence - based resequencing reads .
ESPRIT - Forest	Application	0.92	train	In this paper we developed a new algorithm called ESPRIT - Forest for parallel hierarchical clustering of sequences .

Retrieval Examples: Sentence Similarity

split	sim	text
test	1.00	To this end , we have developed an automated web - based tool , PhosphOrtholog , which allows batch processing and mapping of large species - specific PTM datasets to compare overlap at a site - specific level .
train	0.99	Here we present a tool , Podbat (Positioning database and analysis tool) , that incorporates data from various sources and allows detailed dissection of the entire range of chromatin modifications simultaneously .
train	0.99	We designed and developed a new method , MSACompro , to synergistically incorporate predicted secondary structure , relative solvent accessibility , and residue - residue contact information into the currently most accurate posterior probability - based MSA methods to improve the accuracy of mult... specifically designed to compress MAF (Multiple Alignment Format) files .

NER Recall Confusion



RAG Performance for Task 1 (NER)

Paradigm	Retrieval	F1	Model	parameter
Finetuned	-	0.599	SciBERT	-
Prompt	Random	0.483	GPT 3-5	Random k=7
Prompt	Random	0.525	GPT 3-5	Random all entity types shown
Prompt	Sim. sentences	0.647	GPT 3.5	topn=10
Prompt	Sim. entities	0.624	GPT 3.5	topn=7 x n entities
Prompt	Random	0.574	GPT 4	Random all entity types shown
Prompt	Sim. sentences	0.677	GPT 4	topn=10
Prompt	Sim. entities	0.679	GPT 4	topn=7 x n entities

- #1 System for **SOMD** Subtask 2 and 3
- Entity similarity catch similar entities **and** similar context
- Best on NER for low data regime
- Single-Choice QA for RE: best performing Approach

SOMD Results + RE Baselines

Task #	User	F1	Precision	Recall	Task #	User	F1	Precision	Recall	
1	1	.740	.761	.750	2	1	ours	.838	.835	.847
1	2	david-s477	.692	.739	2	2	phinx	.743	.745	.748
1	3	ThuyNT03	.678	.729	3	1	ours	.916	.911	.924
1	4	ours	.652	.679	3	2	phinx	.897	.900	.897
1	5	vampire	.648	.682	3	-	baseline	.864	.857	.875
					3	-	necessary	.933		.415